

FIG. 1

DEST AVAILABLE COUT

[SEQ. ID NO: 3]

X-C-C-T-T-G-A-G-A-T-T-T-C-C-C-T-C
5'

G-G-A-A-C-T-C-T-A-A-G-G-G-A-G-X
[SEQ. ID NO: 4]

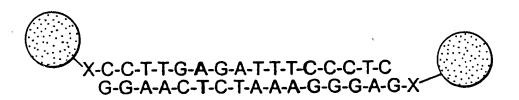




FIG. 2

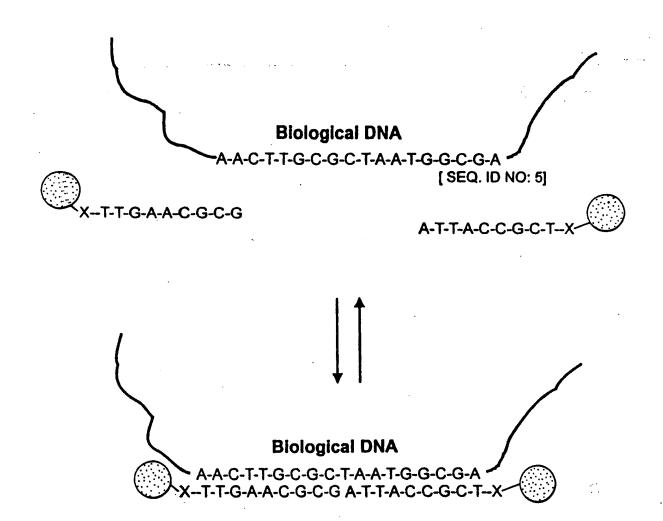
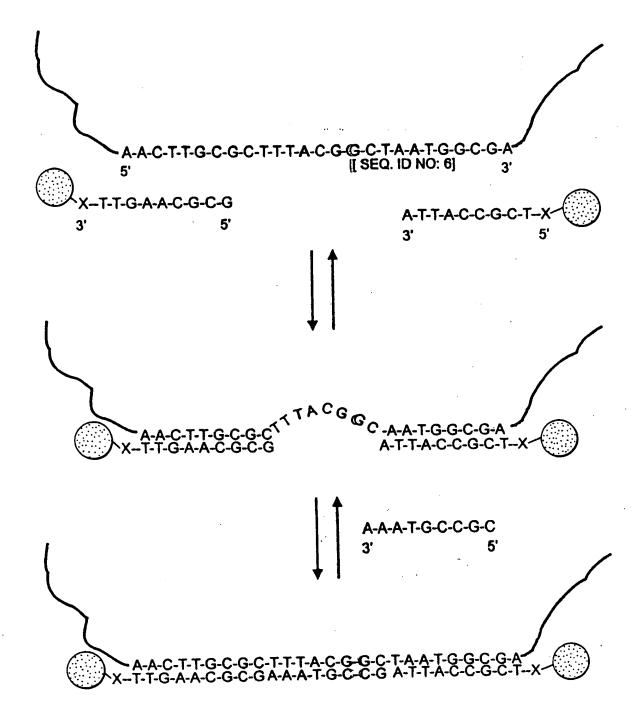
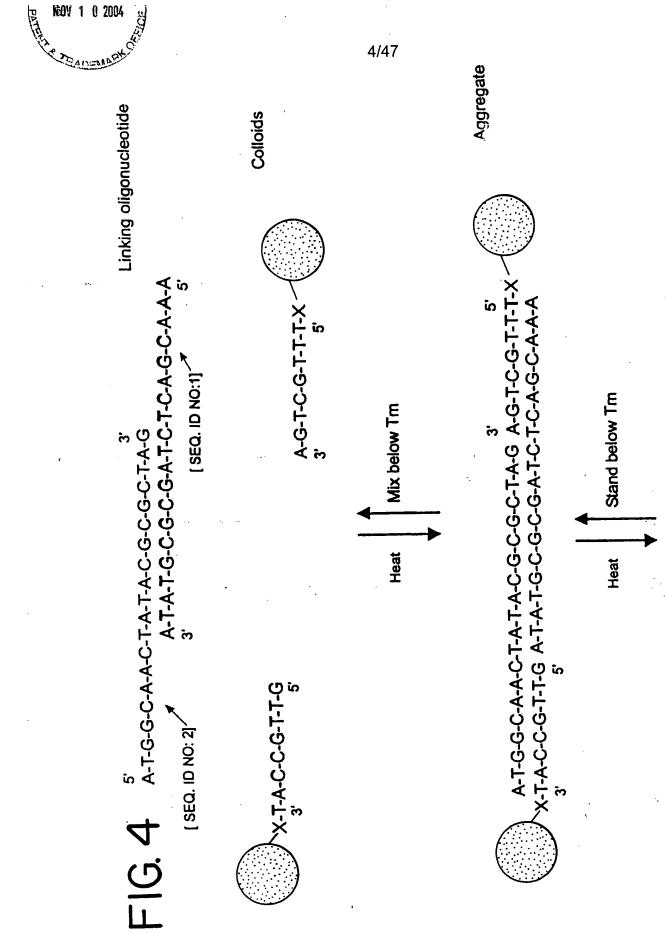




FIG. 3





Precipitate (formed by further cross-linking)



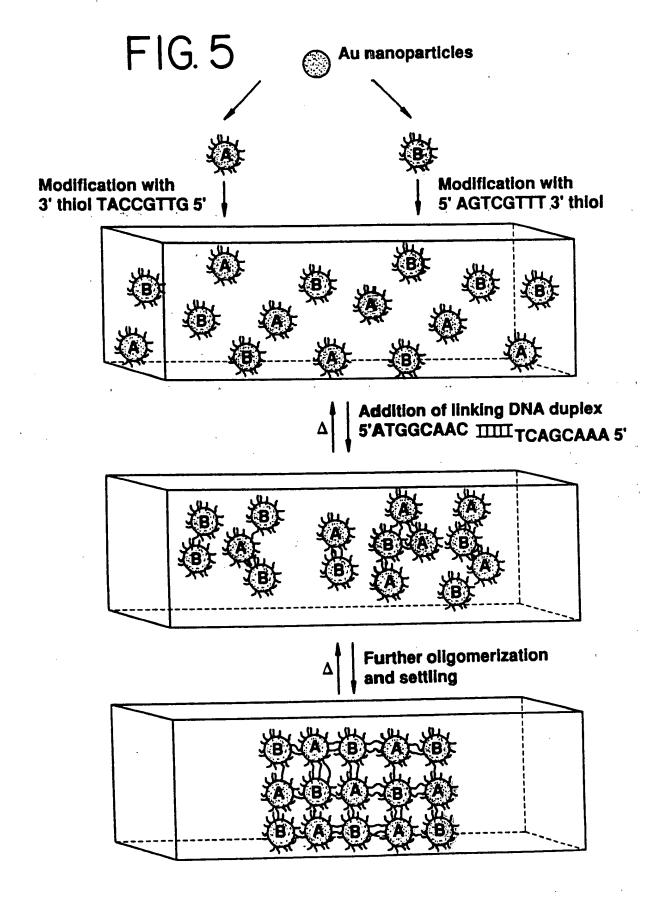
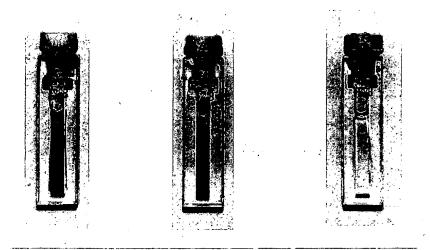


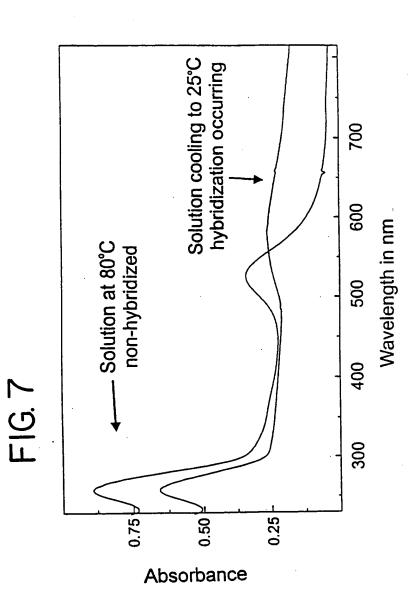


FIG. 6A FIG. 6B FIG. 6C



The second second





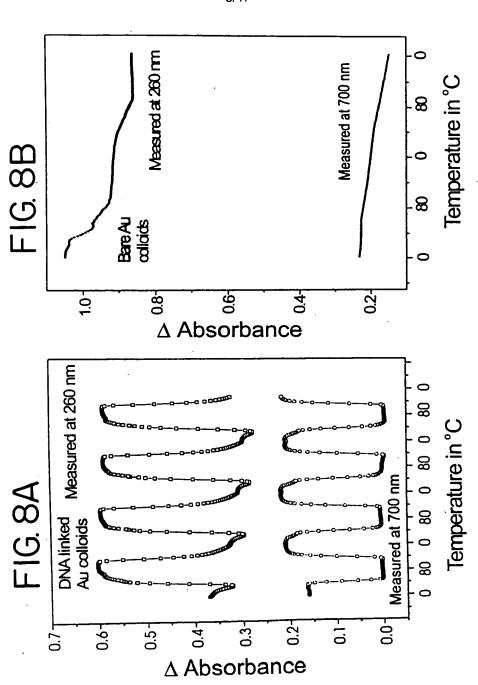




FIG. 9A

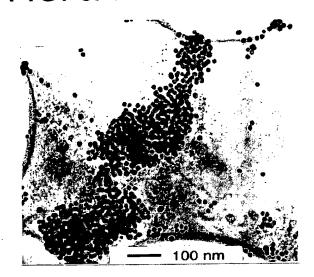


FIG. 9B

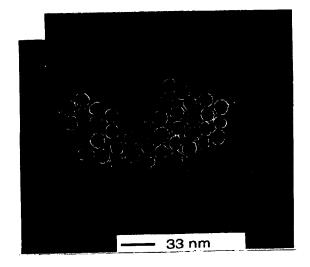




FIG. 10

[SEQ ID NO: 7]

5'
3'
X-T-C-T-C-C-T-T-C-C-C-T-T-T-C
A-G-A-G-G-A-A-G-G-X
3'
[SEQ ID NO: 8]

3, T-C-T-C-C-T-T-C-C-C-T-T-T-T-C 5' [SEQ ID NO: 9]



5' X-T-C-T-C-C-T-T-C-C-C-T-T-T-C A-G-A-G-G-A-A-A-A-G-X T-C-T-C-C-T-T-C-C-C-T-T-T-C-5'

_[SEQ. ID NO: 10]

T-A-C-C-G-T-T-G-A-T-A-T-G-C-G-A-T-C-T-C-A-G-C-A-A-S-

A decimal of page 1

SPE NOV: 1 0 2004 PACIFICAL PROPERTY 12/47 FIG. 12A Complementary Target [SEQ. ID NO: 14] [SEQ. ID NO:12] 1 5' A-G-C-A-T-G-G-T-C-G-A-T-A-G-G-A-A-A-C-G-A-C-T-C-T-A-G-C-G-C [SEQ. ID NO:13] **Probes without Target** 3' T-C-G-T-A-C-C-A-G-C-T-A-T-C-C T-T-T-G-C-T-G-A-G-A-T-C-G-C-G Half Complementary Target 5' A-G-C-A-T-G-G-T-C-G-A-T-A-G-G-A[T-G-G-C]-A[A-C-T-A-T-A]C-G-C SEQ. ID NO: 15 FIG. 12D Target - 6 bp 3' T-C-G-T-A-C-C-A-G-C-T-A-T-C-C T-T-T-G-C-T-G-A-G-A-T-C-G-C-G-5' G-T-C-G-A-T-A-G-G-A-A-A-C-G-A-C-T-C-T-A-G-C-G-C >[SEQ. ID NO:16] One bp Mismatch [SEQ. ID NO: 17] Two bp Mismatch

[SEQ. ID NO:18]



FIG. 13A

transparent substrate

Modified DNA chemisorbed onto solld substrate

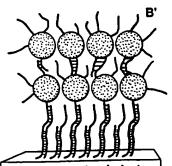
Analyte DNA

transparent substrate

Analyte DNA hybridized onto substrate

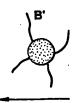
DNA modified colloids

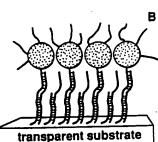




transparent substrate

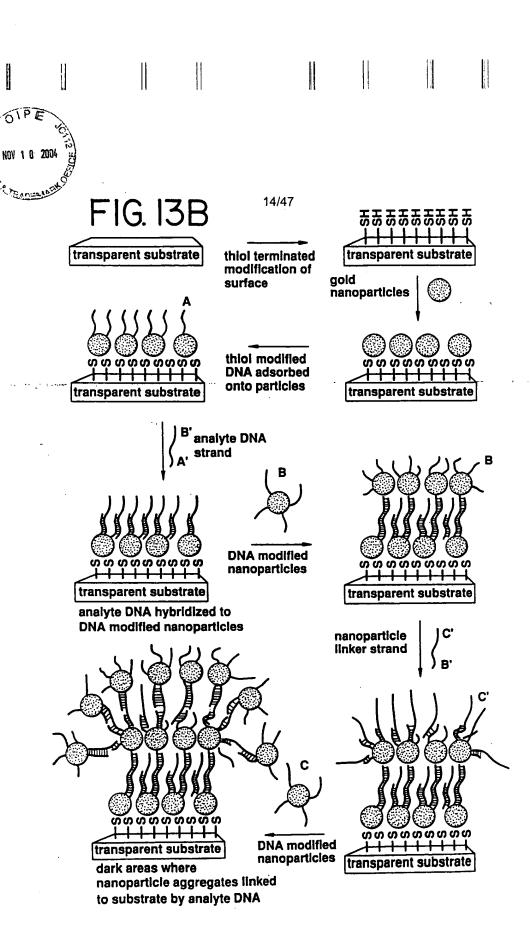
Dark areas where nanoparticle aggregates are linked to substrate surface by analyte DNA





transparent substrate

DNA modified colloids hybridized to bound analyte DNA



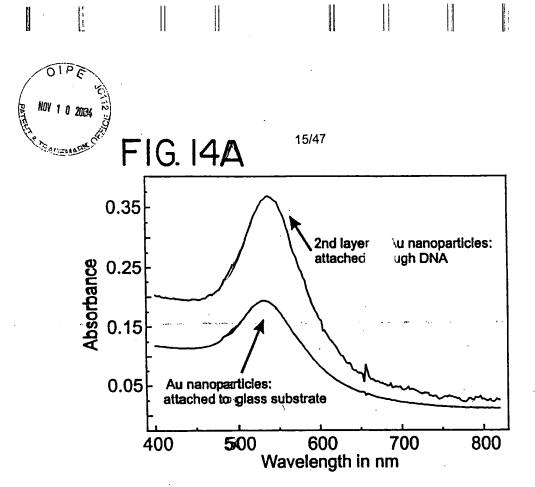
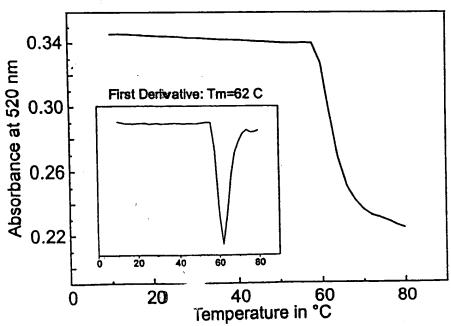
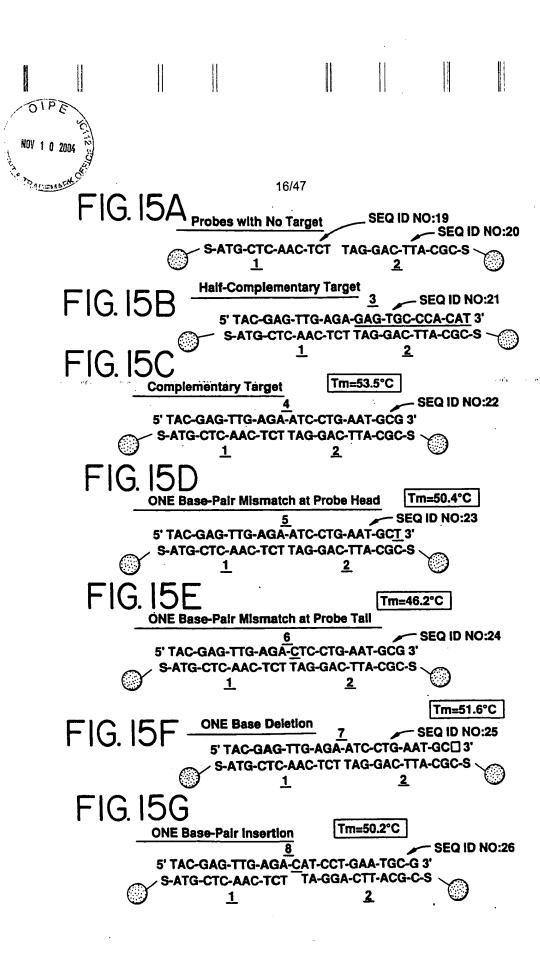


FIG. 14B







24 Base Template

FIG. 16A

5' TAC-GAG-TTG-AGA-ATC-CTG-AAT-GCG 3'

S-ATG-CTC-AAC-TCT TAG-GAC-TTA-CGC-S > 1

FIG. 16B

48 Base Template with Complementary 24 Base Filler

5' TAC-GAG-TTG-AGA-CCG-TTA-AGA-CGA-GGC-AAT-CAT-GCA-ATC-CTG-AAT-GCG 3' -> S-ATG-CTC-AAC-TCT GGC-AAT-TCT-GCT-CCG-TTA-GTA-CGT TAG-GAC-TTA-CGC-S

FIG. 16C

72 Base Template with Complementary 48 Base Filler

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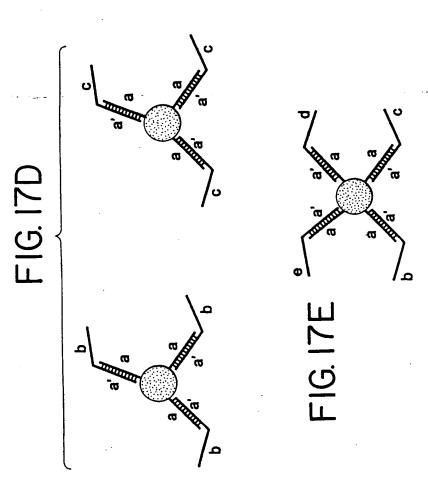
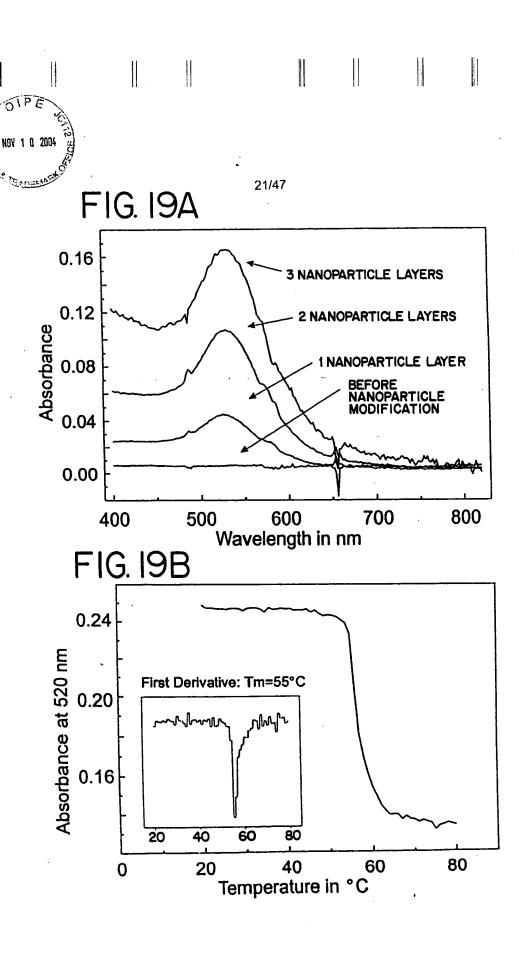
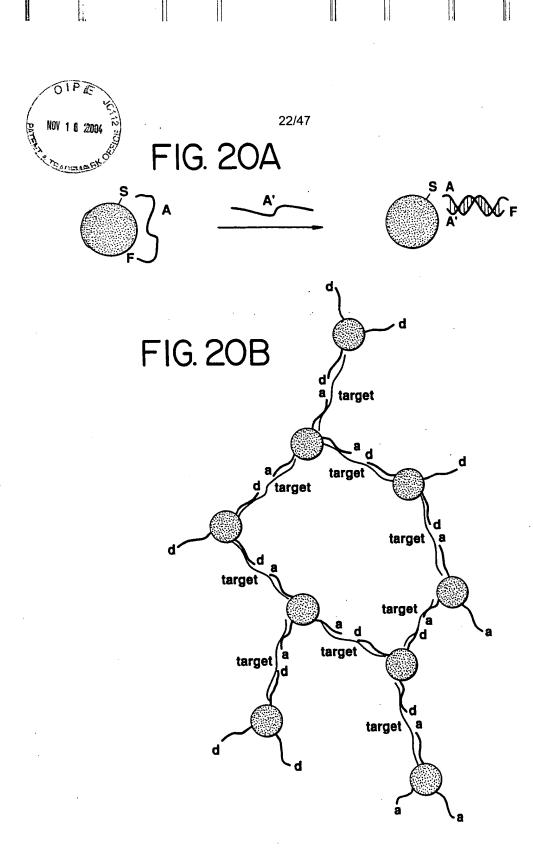


FIG. 18

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Red

Excess Au probes

pass through

All Au probes pass through membrane

membrane

F1G. 21

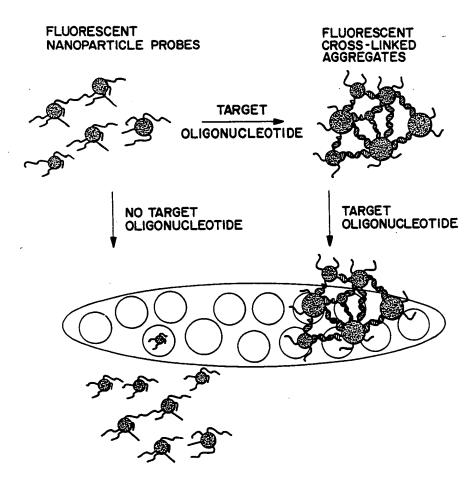
nanoparticle probes

Oligonucleotide modified Au

* MADEMAR

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FIG. 22



THE FLUORESCENT NANOPARTICLE PROBES PASS THROUGH THE MEMBRANE

THE FLUORESCENT CROSS-LINKED AGGREGATES ARE RETAINED BY THE MEMBRANE



FIG. 23

Anthrax PCR Product

5'G GCG GAT GAG TCA GTA GTT AAG GAG GCT CAT AGA GAA GTA ATT AAT 3'C CGC CTA CTC AGT CAT CAA TTC CTC CGA GTA TCT CTT CAT TAA TTA

TCG TCA ACA GAG GGA TTA TTG TTA AAT ATT GAT AAG GAT ATA AGA AAA AGC AGT TGT CTC CCT AAT AAC AAT TTA TAA CTA TTC CTA TAT TCT TTT

ATA TTA TOO AGG GTT ATA TTG TAG AAA TTG AAG ATA CTG AAG GGC TT 3' TAT AAT AGG TCC CAA TAT AAC ATC TTT AAC TTC TAT GAC TTC CCG AA 5'

141 mer Anthrax PCR product [SEQ ID NO:36]

3' CTC CCT AAT AAC AAT

[SEQ ID NO:37]

3' TTA TAA CTA TTC CTA

[SEQ ID NO:38]

Oligonucleotide-Nanoparticle Probes

Blocker Oligonucleotides

3' C CGC CTA CTC AGT CAT CAA TTC CTC CGA GT

3' A TCT CTT CAT TAA TTA AGC AGT TGT

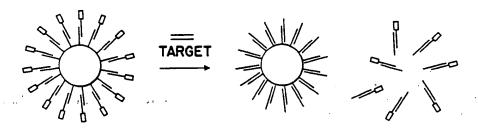
3' TAT TCT TTT TAT AAT AGG TCC CAA TAT

3' AAC ATC TTT AAC TTC TAT GAC TTC CCG AA

[SEQ ID NO:41]



FIG. 24



SATELLITE PROBE

DETECTION SIGNAL

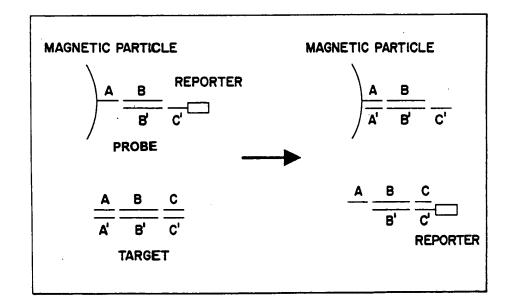
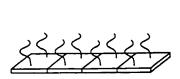




FIG. 25A 1. \(\square \) (TARGET)





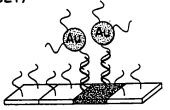
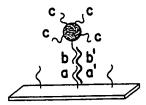


FIG. 25B

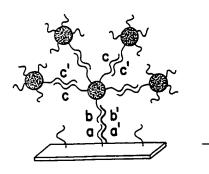












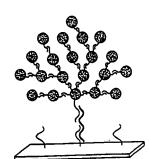




FIG. 26A

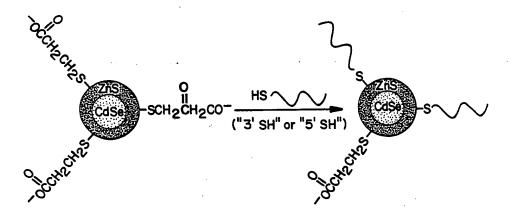
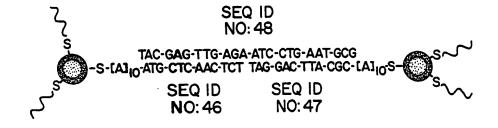
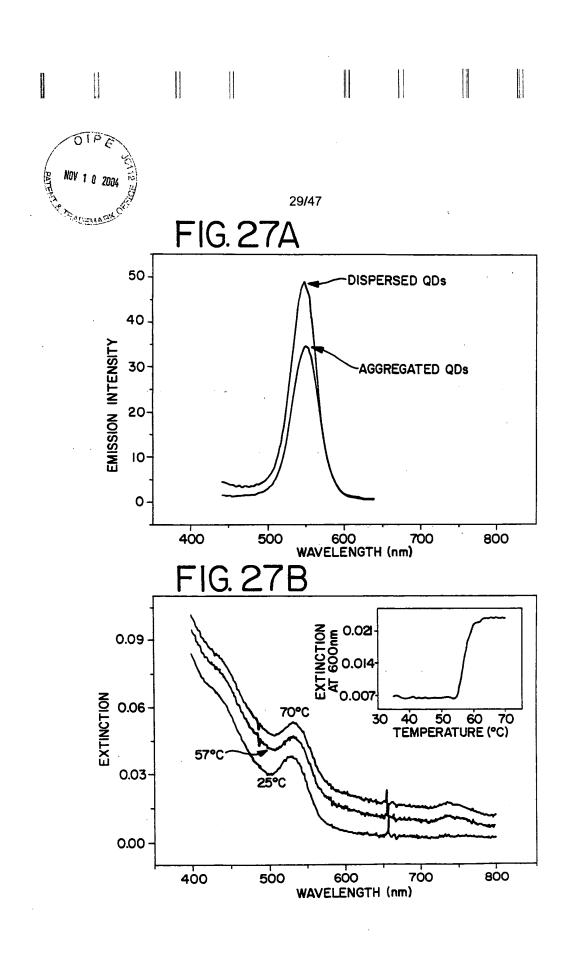


FIG. 26B







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FIG. 27C

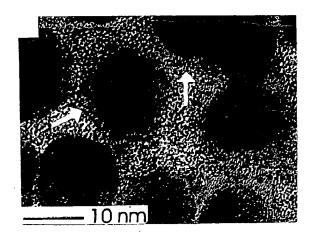
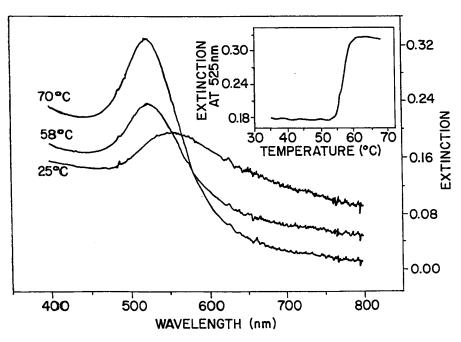


FIG. 27D





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FIG. 28A



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FIG. 28B

FIG. 28C

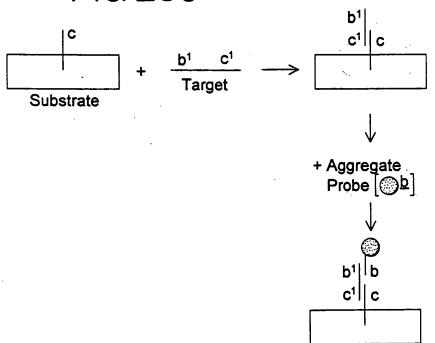




FIG. 28D

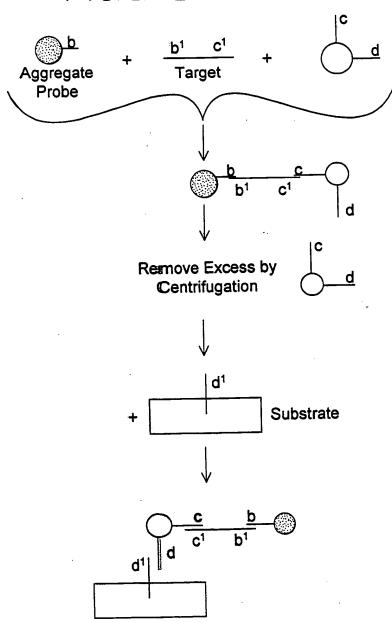
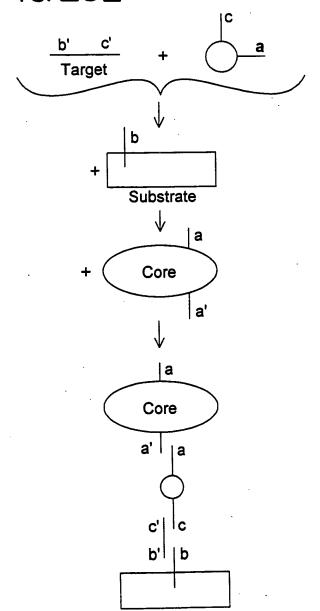
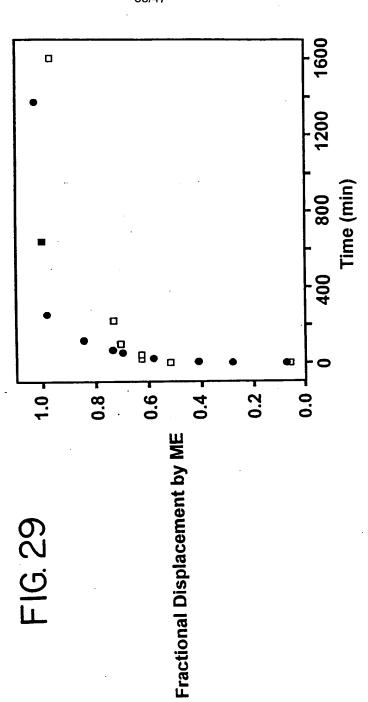
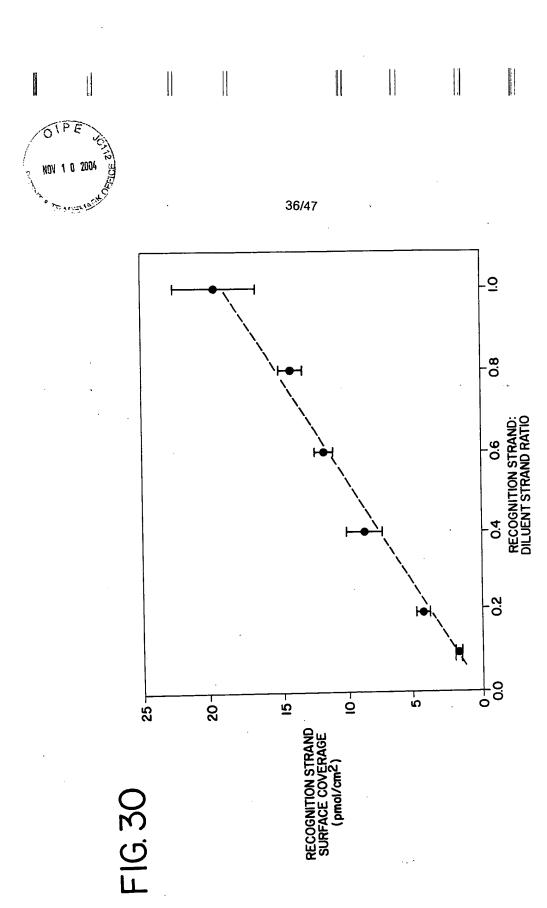


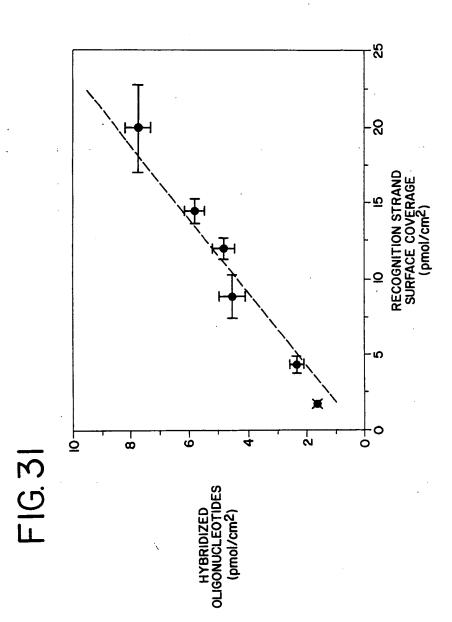


FIG. 28E











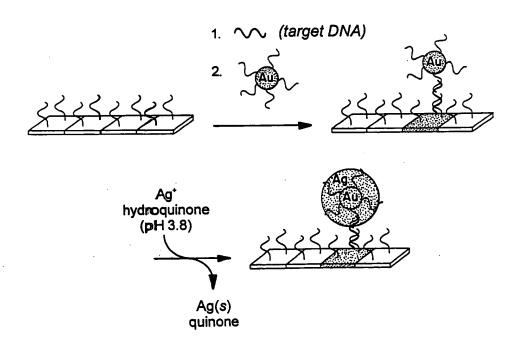
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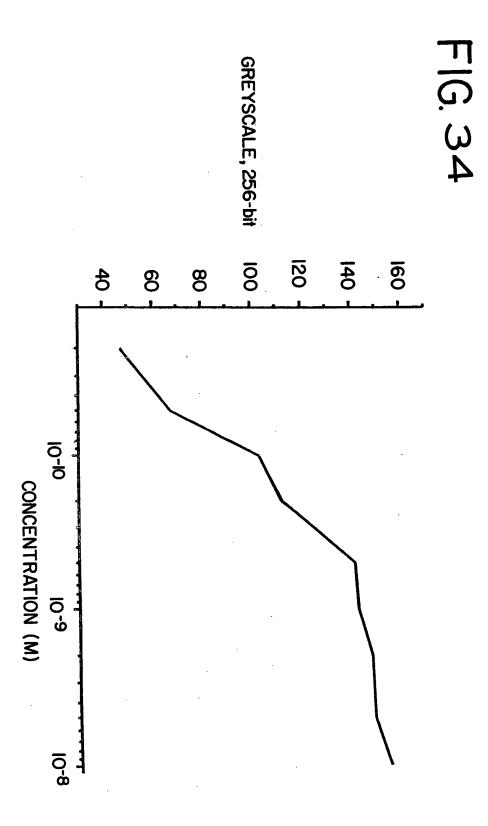
FIG. 32

[SEQIDNO:56]

5' GGA THA TTG TTA--AAT ATT GAT AAG GAT 3' CCT AINT AAC AAT TTA TAA CTA TTC CTA [SEQ: ID NO: 57] [SEQ: ID NO: 58]

> **N** = A (complementary), G,C,T (mismatched)



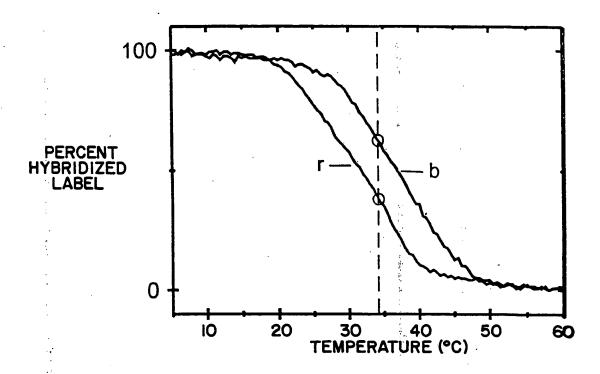


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FIG.35A



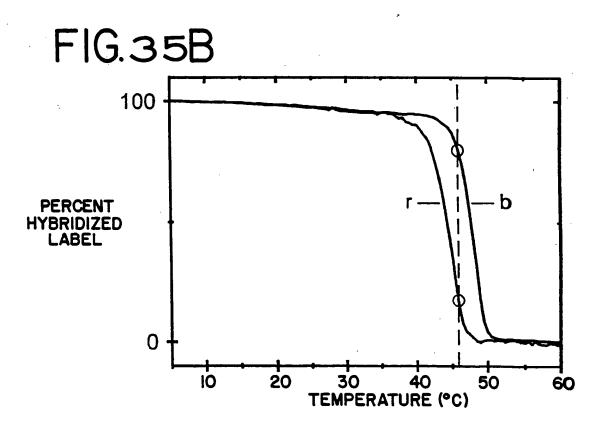




FIG.37A

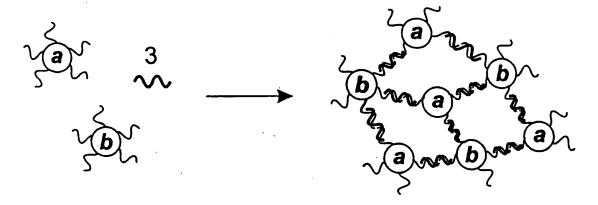
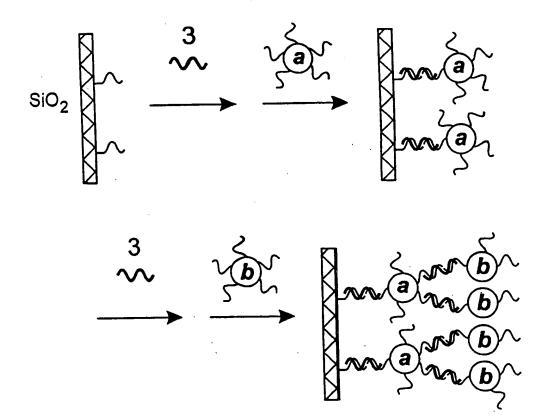
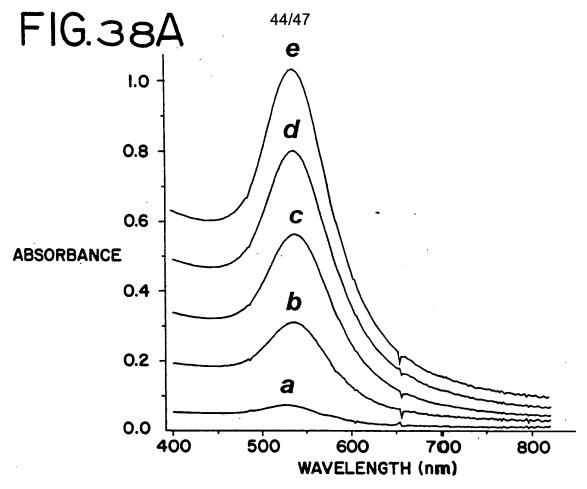
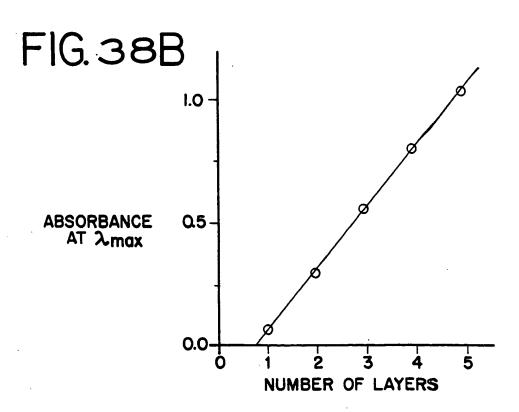


FIG.37B

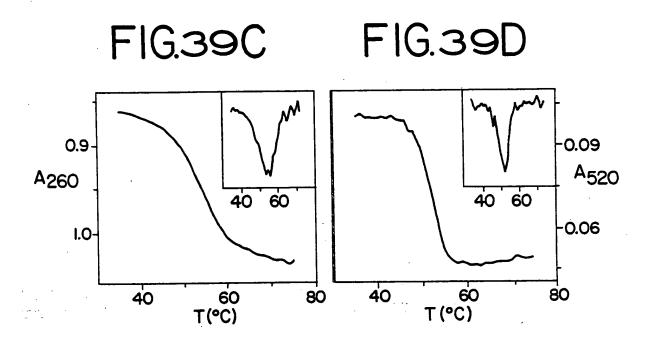


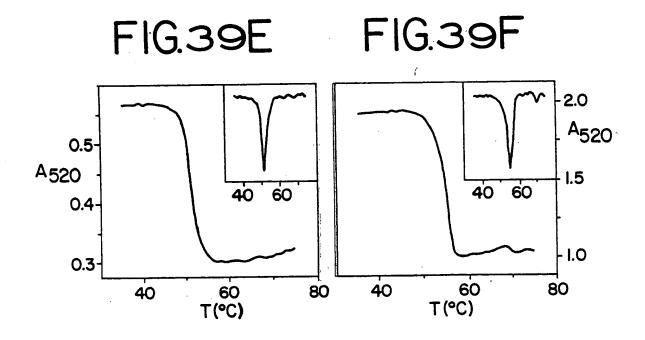














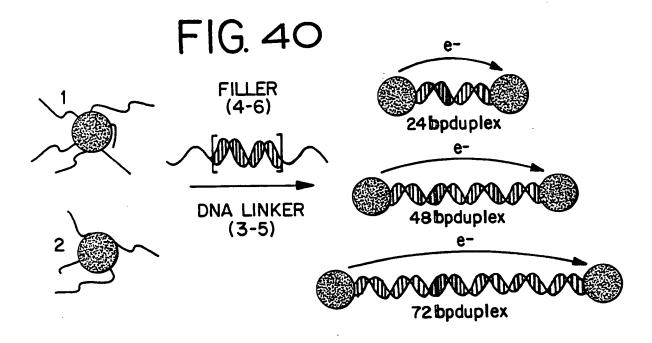


FIG. 4 I



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FIG. 33			
A	B	C	
		II	
D	E	F	G

L**t**/9t

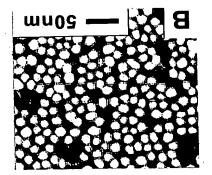


FIG 39B

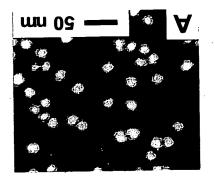


FIG. 39A



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FIG.36A

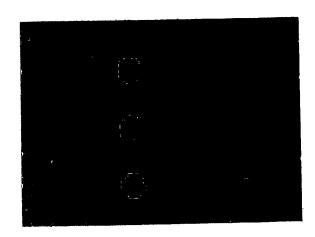
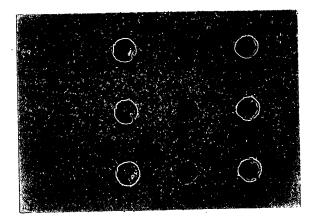


FIG.36B



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